#### SEQUENCE LISTING

#### (1) GENERAL INFORMATION:

- (i) APPLICANT: (US only) BROADY, K. W. and TSENG, A. P. S. (other than US) Analytica, Ltd
- (ii) TITLE OF INVENTION: A METHOD OF TREATMENT
- (iii) NUMBER OF SEQUENCES: 45
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: DAVIES COLLISON CAVE
  - (B) STREET: 1 LITTLE COLLINS STREET
  - (C) CITY: MELBOURNE
  - (D) STATE: VICTORIA
  - (E) COUNTRY: AUSTRALIA
  - (F) ZIP: 3000
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: International PCT
  - (B) FILING DATE: 1999/11/12
- (vii) PRIORITY DATA:
  - (A) APPLICATION NUMBER: 60/108254
  - (B) FILING DATE: 1998/11/12
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: HUGHES, DR E JOHN L
  - (ix) TELECOMMUNICATION INFORMATION:
    - (A) TELEPHONE: +61 3 9254 2777
    - (B) TELEFAX: +61 3 9254 2770
    - (C) TELEX: AA 31787

l	2	) INFORMATION	FOR	SEQ	ID	NO:1:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 202 amino acids
  - (B) TYPE: amino acid
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

Met Lys Ser Leu Gln Ile Ile Cys Leu Leu Phe Val Leu Val Ala Arg

1 5 10 15

Gly Ser Cys His Ser Cys Glu Ile Cys His Asn Leu Gly Arg Asp Cys
20 25 30

Glu Thr Glu Glu Ala Glu Glu Cys Ala Ser Pro Glu Asp Gln Cys Gly
35 40 45

Thr Val Leu Met Glu Val Ser Ser Ala Pro Ile Ser Phe Arg Ser Ile
50 55 60

His Arg Asn Cys Phe Ser Ser Ser Leu Cys Lys Leu Glu Arg Phe Asp 65 70 75 80

Ile Asn Ile Gly His Asp Ser Tyr Leu Arg Gly Arg Ile His Cys Cys
85 90 95

Asp Glu Ala Arg Cys Glu Ala Gln Gln Phe Pro Gly Leu Pro Leu Ser 100 105 110

Phe Pro Asn Gly Tyr His Cys Pro Gly Ile Leu Gly Val Phe Ser Val 115 120 125

Asp Ser Ser Glu His Glu Ala Ile Cys Arg Gly Thr Glu Thr Lys Cys 130 135. 140

Ile Asn Leu Ala Gly Phe Arg Lys Glu Arg Phe Pro Gly Asp Ile Ala 145 150 155 160

Tyr Asn Ile Lys Gly Cys Thr Ser Ser Cys Pro Glu Leu Arg Leu Ser 165 170 175

Asn Arg Thr His Glu Glu Asp Arg Asn Gly Leu Ile Lys Val Glu Cys 180 185 190

ü

Thr Asp Ala Ser Lys Ile Thr Pro Ser Glu
195 200

- (2) INFORMATION FOR SEQ ID NO:2:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 202 amino acids
    - (B) TYPE: amino acid
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: protein
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met Ile Ser Leu Gln Ile Ile Cys Phe Leu Phe Val Leu Val Ala Arg

1 5 10 15

Gly Ser Cys His Ser Cys Glu Ile Cys Arg Asn Phe Gly Lys Asp Cys
20 25 30

Glu Ser Glu Glu Ala Glu Glu Cys Ala Ser Pro Glu Asp Gln Cys Gly
35 40 45

Thr Val Leu Leu Glu Ile Ser Ser Ala Pro Ile Ser Phe Arg Ser Ile
50 55 60

His Arg Asn Cys Phe Ser Ser Ser Leu Cys Lys Leu Glu His Phe Asp
65 70 75 80

Ile Asn Ile Gly His Asp Ser Tyr Val Arg Gly Arg Ile His Cys Cys 85 90 95

Asp Glu Glu Arg Cys Glu Ala Gln Gln Phe Pro Gly Leu Pro Pro Ser

Leu Pro Asn Gly Tyr His Cys Pro Gly Ile Leu Gly Ala Phe Ser Val

Asp Ser Ser Glu His Glu Ala Ile Cys Arg Gly Thr Glu Thr Lys Cys
130 140

Ile Asn Leu Ala Gly Phe Arg Lys Glu Arg Tyr Pro Val Asp Ile Ala 145 150 155 160

Tyr Asn Ile Thr Gly Cys Thr Ser Ser Cys Pro Glu Leu Lys Leu Ser 165 170 175

Asn	Arg	Thr	His	Ala	Glu	Ang	Arg	Asn	Ala	Leu	Ile	Thr	Leu	Asp	Суз
	•		180					185					190		

Thr Asp Ala Ser Lys Ile Ala Pro Ser Glu
195 200

#### (2) INFORMATION FOR SEQ ID NO:3:

#### (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 609 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

#### (ii) MOLECULE TYPE: CDNA

#### (ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1..606

#### (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

ATG AAA TCC CTA CAG ATC ATC TGT CCT CTT TTC GTT TTG GTA GCC AGA

Met Lys Ser Leu Gln Ile Ile Cys Pro Leu Phe Val Leu Val Ala Arg

1 5 10 15

GGA AGC TGT CGC TCA TGT GAA ATT TGT CAC AAT TTT GGA AAA GAT TGC 9.

Gly Ser Cys Arg Ser Cys Glu Ile Cys His Asn Phe Gly Lys Asp Cys

20 25 30

GAG AGT GAG GAG GCA GAG GAA TGT GCC TCT CCA GAA GAT CAA TGT GGC

144
Glu Ser Glu Glu Ala Glu Glu Cys Ala Ser Pro Glu Asp Gln Cys Gly

35

40

45

ACA GTG TTG CTG GAG ATT TCA TCA GCA CCT ATT TCC TTC CGA TCC ATT

192
Thr Val Leu Leu Glu Ile Ser Ser Ala Pro Ile Ser Phe Arg Ser Ile

50 55 60

CAT AGG AAC TGT TTC TCA TCC AGC CTC TGC AAA CTT GAA CAC TTT GAT

His Arg Asn Cys Phe Ser Ser Leu Cys Lys Leu Glu His Phe Asp

70

75
80

ATA AAT ATT GGA CAT GAT TCC TAT GTG AGA GGA AGA ATC CAC TGT TGT

11e Asn Ile Gly His Asp Ser Tyr Val Arg Gly Arg Ile His Cys Cys

85

90
95

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			193					200										

## (2) INFORMATION FOR SEQ ID NO:4:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 28 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

Leu Glu Cys Glu Ile Cys Ile Gly Leu Gly Leu Glu Cys Asn Thr Xaa 1 5 10 15 Thr Lys Thr Cys Asp Ala Asn Gln Asp Thr Cys Val 20 25

- (2) INFORMATION FOR SEQ ID NO:5:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 14 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Leu Glu Cys Glu Ile Cys Ile Gly Leu Gly Leu Glu Cys Asn

- (2) INFORMATION FOR SEQ ID NO:6:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 5 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

Ala Leu Ser Tyr Lys

- (2) INFORMATION FOR SEQ ID NO:7:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

Ser Cys Gly Thr Ser Asp Thr Cys His Leu Asn Tyr Val Glu Thr Thr 1 5 10 15

Pro His Asn

- (2) INFORMATION FOR SEQ ID NO:8:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 18 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

Thr Cys Asp Ala Asn Gln Asp Thr Cys Val Thr Phe Gln Thr Glu Val

Ile Arg

- (2) INFORMATION FOR SEQ ID NO:9:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 8 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

Ala Pro Val Thr Leu Gly Leu Ile
1 5

- (2) INFORMATION FOR SEQ ID NO:10:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 10 amino acids
    - (B) TYPE: amino acid

- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

Glu Cys Thr Glu His Leu Val Ser Cys Arg 1 5 10

- (2) INFORMATION FOR SEQ ID NO:11:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 13 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

Phe Trp Asn Val Leu Glu Asp Val Glu Val Asp Phe Lys
1 5 10

- (2) INFORMATION FOR SEQ ID NO:12:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 29 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

His Ser Cys Glu Ile Cys His Asn Phe Gly Arg Asp Cys Gln Ser Asp 1 5 10 15

Glu Ala Glu Glu Cys Ala Ser Pro Glu Asp Gln Cys Gly

20

25

- (2) INFORMATION FOR SEQ ID NO:13:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 29 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

His Ser Cys Glu Ile Cys His Asn Leu Gly Lys Asp Cys Glu Thr Glu

Glu Thr Glu Glu Cys Ala Ser Pro Glu Asp Gln Cys Gly

- (2) INFORMATION FOR SEQ ID NO:14:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 5 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

Ile Thr Pro Ser Glu 5

(2) INFORMATION FOR SEQ ID NO:15:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 6 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

Arg Phe Asp Ile Asn Ile

- (2) INFORMATION FOR SEQ ID NO:16:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 6 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

Ile Asn Leu Ala Gly Phe

- (2) INFORMATION FOR SEQ ID NO:17:
  - (i) SEOUENCE CHARACTERISTICS:
    - (A) LENGTH: 8 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

Ala Ser Lys Ile Thr Pro Ser Glu 1 5

- (2) INFORMATION FOR SEQ ID NO:18:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 7 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Tyr Pro Gly Asp Ile Ala Ile

- (2) INFORMATION FOR SEQ ID NO:19:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 28 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Leu Glu Cys Glu Ile Cys Ile Gly Leu Gly Leu Glu Cys Asn Thr Trp

1 5 10 15

Thr Lys Thr Cys Asp Ala Asn Gln Asp Thr Cys Val

- (2) INFORMATION FOR SEQ ID NO:20:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 5 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

Ala Leu Ser Tyr Lys

- (2) INFORMATION FOR SEQ ID NO:21:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19 amino acids
      - (B) TYPE: amino acid
      - (C) STRANDEDNESS: single
      - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

Ser Cys Gly Thr Ser Asp Thr Cys His Leu Asn Tyr Val Glu Thr Thr

1 10 15

Pro His Asn

(2) INFORMATION FOR SEQ ID NO:22:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 18 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

Thr Cys Asp Ala Asn Gln Asp Thr Cys Val Thr Phe Gln Thr Glu Val

1 5 10 15

Ile Arg

- (2) INFORMATION FOR SEQ ID NO:23:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 8 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

Ala Pro Val Thr Leu Gly Leu Ile
1 5

- (2) INFORMATION FOR SEQ ID NO:24:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 10 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single

- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

Glu Cys Thr Glu His Leu Val Ser Cys Arg

- (2) INFORMATION FOR SEQ ID NO:25:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 13 amino acids
    - (B) TYPE: amino acid
      - (C) STRANDEDNESS: single
      - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

Phe Trp Asn Val Leu Glu Asp Val Glu Val Asp Phe Lys

1 5 10

- (2) INFORMATION FOR SEQ ID NO:26:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 10 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

Gly Ser Glu Asn Gln Cys Lys Ser Ile Ile 1 5 . 10

- (2) INFORMATION FOR SEQ ID NO:27:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 22 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single.
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

Val Asn Pro Pro Asn Gly Leu Gln Cys Pro Gly Cys Leu Gly Leu Ser

1 10 15

Ser Leu Glu Cys Thr Glu 20

- (2) INFORMATION FOR SEQ ID NO:28:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 14 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

Cys Gly Thr Ser Asp Thr Cys His Leu Asn Tyr Val Glu Thr

1 5 10

- (2) INFORMATION FOR SEQ'ID NO:29:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 7 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:

Glu Phe Gly Leu Phe Phe Arg

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- (2) INFORMATION FOR SEQ ID NO:30:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 183 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

His Ser Cys Glu Ile Cys His Asn Phe Gly Lys Asp Cys Glu Gly Gly

1 5 10 15

Glu Thr Glu Glu Cys Ala Ser Pro Glu Asp Gln Cys Gly Thr Val Leu 20 25 30

Met Glu Val Ser Thr Ala Pro Ile Ser Phe Arg Ser Ile His Arg Asn 35 40 45

Cys Phe Ser Ser Leu Cys Lys Leu Glu Arg Phe Asp Ile Asn Ile

50

55

60

Gly His Asp Ser Phe Leu Arg Gly Arg Ile His Cys Cys Asp Glu Ala 65 70 75 80

Arg Cys Glu Ala Gln Gln Phe Pro Gly Leu Pro Leu Ser Phe Pro Asn 85 90 95

Gly Tyr His Cys Pro Gly Ile Leu Gly Leu Phe Ser Val Asp Ser Ser 100 105 110

Glu His Glu Ala Ile Cys Arg Gly Thr Glu Thr Lys Cys Ile Asn Leu 115 120 125

Ala Gly Phe Arg Arg Glu Arg Phe Pro Gly Asp Ile Ala Tyr Asn Ile 130 135 140

Lys Gly Cys Thr Ser Ser Cys Pro Glu Leu Arg Leu Ser Asn Arg Thr.

145 150 155 160

His Glu Glu His Arg Asn Asp Leu Ile Lys Val Glu Cys Thr Glu Ala 165 170 175

Ser Lys Asn Thr Pro Ser Glu 180

## (2) INFORMATION FOR SEQ ID NO:31:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 182 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide

#### (xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:

His Ser Cys Glu Ile Cys His Asn Phe Gly Lys Asp Cys Gln Ser Asp

1	5,	.10		15
			;	

Glu Thr Glu Glu Cys Ala Ser Ala Glu Asp Gln Cys Gly Thr Val Leu 20 25 30

Met Glu Val Ser Ser Ala Pro Ile Ser Phe Arg Ser Ile His Arg Lys
35 40 45

Cys Phe Ser Ser Ser Leu Cys Lys Leu Glu Arg Phe Asp Ile Asn Ile 50 55 60

Gly His Asp Ser Tyr Leu Arg Gly Arg Ile His Cys Cys Asp Glu Ala 65 70 75 80

Arg Cys Glu Ala Gln Gln Phe Pro Gly Leu Pro Leu Ser Phe Pro Asn 85 90 95

Gly Tyr His Cys Pro Gly Ile Leu Gly Val Phe Ser Val Asp Ser Ser 100 105 110

Glu His Glu Ala Ile Cys Arg Gly Thr Glu Thr Lys Cys Ile Asn Leu 115 120 125

Ala Gly Phe Arg Lys Glu Arg Tyr Pro Ile Asp Ile Ala Tyr Asn Ile 130 135 140

Lys Gly Cys Thr Ser Ser Cys Pro Glu Leu Arg Leu Asn Arg Thr His 145 150 155 160

Glu Glu His Arg Asn Asp Leu Ile Lys Val Glu Cys Thr Glu Ala Ser 165 170 175

Lys Ile Thr Pro Ser Glu 180

#### (2) INFORMATION FOR SEQ ID NO: 32:

#### (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 183 amino acids
- (B) TYPE: amino acid.
- (C) STRANDEDNESS: single

- (D) TOPOLOGY; linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:

His Ser Cys Glu Ile Cys His Asn Phe Gly Lys Asp Cys Glu Gly Gly

1 10 15

Val Thr Glu Glu Cys Ala Ser Pro Glu Asp Gln Cys Gly Thr Val Leu 20 25 30

Leu Glu Val Ser Thr Ala Pro Ile Ser Thr Arg Thr Ile His Arg Asn 35 40 45

Cys Phe Ser Ser Ser Leu Cys Lys Leu Glu Arg Phe Asp Ile Asn Ile
50 55 60

Gly His Asp Ser Tyr Met Arg Gly Arg Ile His Cys Cys Asp Glu Ala 65 70 75 80

Arg Cys Glu Ala Gln Gln Phe Pro Gly Leu Pro Leu Ser Phe Pro Asn 85 90 95

Gly Tyr His Cys Pro Gly Ile Leu Gly Leu Phe Ser Val Asp Ser Ser 100 105 110

Glu His Glu Ala Ile Cys Arg Gly Ser Glu Thr Lys Cys Ile Lys Ile 115 120 125

Ala Gly Phe Arg Arg Glu Arg Tyr Pro Ile Asp Ile Ala Tyr Asn Ile 130 135 140

Lys Gly Cys Thr Ser Ser Cys Pro Glu Leu Arg Leu Ser Asn Arg Thr
145 150 155 . 160

His Glu Glu His Arg Asn Asp Leu Ile Lys Val Glu Cys Thr Asp Ala 165 170 175

- 20 -

Ser Lys Ile Thr Pro Ser Glu 180

# (2) INFORMATION FOR SEQ ID NO:33:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 181 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:

Leu Glu Cys Glu Ile Cys Ile Gly Leu Gly Leu Glu Cys Asn Thr Trp

1 5 10 15

Thr Lys Thr Cys Asp Ala Asn Gln Asp Thr Cys Val Thr Phe Gln Thr

Glu Val Ile Arg Ala Pro Val Ser Leu Ser Leu Ile Ser Lys Ser Cys
35 40 45

Gly Thr Ser Asp Thr Cys His Leu Asn Tyr Val Glu Thr Ser Pro His 50 55 60

Asn Glu Leu Thr Val Lys Thr Lys Arg Thr Cys Cys Thr Gly Glu Glu 65 70 75 80

Cys Lys Thr Leu Pro Pro Pro Val Leu Gly His Lys Val Asn Pro Pro 85 90 95

Asn Gly Leu Gln Cys Pro Gly Cys Leu Gly Leu Ser Ser Lys Glu Cys
100 105 110

Thr Glu His Leu Val Ser Cys Arg Gly Ser Glu Asn Gln Cys Leu Ser

Ile	Ile	Gly	ГÀè	Glu	Phe	Gly	Leu	Phe	Phe	Arg	Ala	Leu	Ser	Tyr	Lys
	130					135					140				

Gly Cys Ala Thr Glu Ser Leu Cys Thr Leu Phe Glu Lys Arg Phe Trp 145 150 155 160

Asn Val Leu Glu Asp Val Glu Val Asp Phe Lys Cys Thr Pro Ala Leu 165 170 175

Pro Lys Ser Ser Gln 180

## (2) INFORMATION FOR SEQ ID NO:34:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 501 base pairs
- (B) TYPE: nucleic acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear

#### (ii) MOLECULE TYPE: DNA

# (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:

CACTCATGTG AAATTTGTCA CAATTTTGGA AAAGATTGCG AGGGTGGGGA GACAGAGGAA 60 TGTGCCTCTC CAGAAGATCA ATGTGGCACA GTGTTGATGG AGGTTTCAAC AGCACCTATT 120 TCCTTCCGAT CCATTCATAG GAACTGTTTC TCATCCAGCC TCTGCAAACT TGAACGCTTT 180 GATATAAATA TTGGACATGA TTCCTTTTTG AGAGGAAGAA TCCACTGTTG TGATGAAGCA 240 AGGTGTGAAG CACAGCAATT TCCTGGACTG CCCCTCTCCT TTCCAAATGG ATACCACTGC 300 CCTGGAATTC TTGGTTTATT CTCAGTGGAC AGCTCTGAAC ATGAAGCTAT TTGCAGAGGA 360 ACTGAAACCA AATGCATTAA CCTTGCGGGA TTCAGAAGAG AAAGATTTCC TGGAGACATC 420 GCTTATAATA TCAAAGGTTG CACTTCTTCT TGTCCAGAAC TGAGGTTGAG CAATAGAACT 480

CACGAAGAAC	ATAGAAATGA	G
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501

## (2) INFORMATION FOR SEQ ID NO:35:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 501 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

#### (ii) MOLECULE TYPE: DNA

# (xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:

CACTCATGTG	AAATTTGTCA	CAATTTTGGA	AAAGATTGCC	AGAGTGACGA	GACAGAGGAA	6
TGTGCCTCTG	CAGAAGATCA	ATGTGGCACG	GTGTTGATGG	AGGTTTCATC	AGCACCTATT	120
TCCTTCCGAT	CCATTCATAG	GAAGTGTTTC	TCATCCAGCC	TCTGCAAACT	TGAACGCTTT	180
GATATAAATA	TTGGACATGA	TTCCTATTTG	AGAGGAAGAA	TCCACTGTTG	TGATGAAGCA	240
AGGTGTGAAG	CACAGCAATT	TCCTGGACTG	CCCCTCTCCT	TTCCAAATGG	ATACCACTGC	300
CCTGGCATTC	TTGGTGTATT	CTCAGTGGAC	AGCTCTGAAC	ATGAAGCTAT	TTGCAGAGGA	360
ACTGAAACCA	AATGCATTAA	CCTTGCGGGA	TTCAGAAAAG	AAAGATATCC	TATAGACATC	420
GCTTATAATA	TCAAAGGTTG	CACTTCTTCT	TGTCCAGAAC	TGAGGTTGAA	TAGAACTCAC	480
GAAGAACATA	GAAATGATCT	A				501

#### (2) INFORMATION FOR SEQ ID NO:36:

#### (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 501 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE; DNA

(vi)	SECTIENCE	DESCRIPTION:	CEA	TD	NO. 1	٠.
127	SECUENCE	DESCRIPTION:	SEC	111	NO.	46.

CACTCATGTG	AAATTTGTCA	CAATTTTGGA	AAAGATTGCG	AGGGTGGGGT	GACAGAGGAA	60
TGTGCCTCTC	CAGAAGATCA	ATGTGGCACA	GTGTTGCTGG	AGGTTTCAAC	AGCACCTATT	120
TCCACCCGAA	CCATTCATAG	GAACTGTTTC	TCATCCAGCC	TCTGCAAACT	TGAACGCTTT	180
GATATAAATA	TTGGACATGA	TTCCTATATG	AGAGGAAGAA	TCCACTGTTG	TGATGAAGCA	240
AGGTGTGAAG	CACAGCAATT	TCCTGGACTG	CCCCTCTCCT	TTCCAAATGG	ATACCACTGC	300
CCTGGCATTC	TTGGTTTATT	CTCAGTGGAC	AGCTCTGAAC	ATGAAGCTAT	TTGCAGAGGA	360
AGTGAAACCA	AATGCATTAA	AATTGCGGGA	TTCAGAAGAG	AAAGATATCC	TATAGACATC	420
GCTTATAATA	TCAAAGGTTG	CACTTCTTCT	TGTCCAGAAC	TGAGGTTGAG	CAATAGAACT	480
CACGAAGAAC	ATAGAAATGA	T				501

#### (2) INFORMATION FOR SEQ ID NO:37:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 825 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

## (ii) MOLECULE TYPE: DNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:

CTTGAGTGTG	AGATTTGTAT	CGGGCTGGGC	CTGGAATGTA	ACACCTGGAC	GAAAACCTGT	60
				,		
GATGCTAATC	AAGATACTTG	TGTTACCTTT	CAAACTGAAG	TGATAAGAGC	CCCTGTGTCC	120

CTCTCTTTGA	TTTCAAAATC	CTGTGGTACT	TCTGACACTT	GCCATCTTAA	CTACGTGGAG	180	
ACGAGTCCAC	ATAATGAACT	' AACAGTGAAG	ACCAAAAGAA	CCTGCTGTAC	TGGGGAGGAA	240	i
TGTAAAACTC	TGCCACCGCC	TGTGCTTGGA	CACAAAGTCA	ACCCACCCAA	CGGACTTCAG	300	
TGTCCTGGAT	GCCTTGGATT	GTCCTCAAAA	GAATGCACTG	AACACCTGGT	TTCCTGCCGG	360	
GGATCTGAAA	ACCAGTGTTT	GTCTATAATT	GGGAAAGAAT	TTGGCCTTTT	CTTCAGAGCA	420	
TTGTCTTATA	AAGGATGTGC	TACGGAGAGT	CTGTGCACTT	TATTTGAGAA	GAGGTTCTGG	480	
AATGTTTTAG	AGGATGTTGA	AACAACATAC	TCCAAAACAG	CCAAAACAGC	CAAAACAGCA	540	
CTACATACTC	CTAACCGTAT	GCACAACAAC	CAAAAATGAA	ATCCCTACAG	ATCATCTGTC	600	
TTCTTTTCGT	TTTGGTAGCC	AGAGGAAGCT	GTCAAAAATG	AAATCCCTAC	AGATCATCTG	·- ·660·	
CTTCTTTTC	GTTTTGGTAG	CCAGAGGAAG	CTGTCAAAAA	TGAAATCCCT	ACAGATCATC	720	
GTCTTCTTT	TCGTTTTGGT	AGCCAGAGGA	AGCTGTACTA	CAACCTAAAT	GAAGTCCCTC	780	
TATTCTGTT	GCCTCTTTGG	CACTTTCTTA	GCTACAGGCA	TGTGT		825	

#### (2) INFORMATION FOR SEQ ID NO:38:

- (i) SEQUENCE CHARACTERISTICS:
  - (A) LENGTH: 19 amino acids
  - (B) TYPE: amino acid
  - (C) STRANDEDNESS: single
  - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

Met Lys Ser Leu Gln Ile Ile Cys Leu Leu Phe Val Leu Val Ala Arg 1 5 10 15 Gly Ser Cys

- (2) INFORMATION FOR SEQ ID NO:39:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:

Met Lys Ser Leu Gin Ile Ile Cys Leu Leu Phe Val Leu Val Ala Arg

1 5 10 15

Gly Ser Cys

- (2) INFORMATION FOR SEQ ID NO:40:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:

Met Lys Ser Leu Gln Ile Ile Cys Leu Leu Phe Val Leu Val Ala Arg

5 10 15

Gly Ser Cys

- (2) INFORMATION FOR SEQ ID NO:41:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 19 amino acids
    - (B) TYPE: amino acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: peptide
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:

Met Lys Ser Leu Leu Phe Cys Cys Leu Phe Gly Thr Phe Leu Ala Thr

1 5 10 15

Gly Met Cys

- (2) INFORMATION FOR SEQ ID NO:42:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 57 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:

ATGAAATCCC TACAGATCAT CTGTCTTCTT TTCGTTTTGG TAGCCAGAGG AAGCTGT

57

- (2) INFORMATION FOR SEQ ID NO:43:
  - (i) SEQUENCE CHARACTERISTICS:

	(A)	LENGTH:	57	base	pairs
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- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: DNA
- (x1) SEQUENCE DESCRIPTION: SEQ ID NO:43:

# ATGAAATCCC TACAGATCAT CTGTCTTCTT TTCGTTTTGG TAGCCAGAGG AAGCTGT

57

- (2) INFORMATION FOR SEQ ID NO:44:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 57 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA
  - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:

## ATGAAATCCC TACAGATCAT CTGTCTTCTT TTCGTTTTGG TAGCCAGAGG AAGCTGT

57

- (2) INFORMATION FOR SEQ ID NO:45:
  - (i) SEQUENCE CHARACTERISTICS:
    - (A) LENGTH: 57 base pairs
    - (B) TYPE: nucleic acid
    - (C) STRANDEDNESS: single
    - (D) TOPOLOGY: linear
  - (ii) MOLECULE TYPE: DNA

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:

ATGAAGTCCC TCTTATTCTG TTGCCTCTTT GGCACTTTCT TAGCTACAGG CATGTGT 57